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CORRECTION

Correction to: Expanding the phenotype of hypomaturation amelogenesis imperfecta due to a novel *SLC24A4* variant

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In the pedigree, one of the individuals was marked as unaffected whereas it is heterozygous for the SLC24A4 mutation. Therefore figure 1 has been corrected, as well as the relevant text in the abstract, the results and in the supplements. The fundamental findings of the study are not affected by this error.

Below is the corrected figure 1.

The online version of the original article can be found at https://doi.org/10.1007/s00784-020-03222-7

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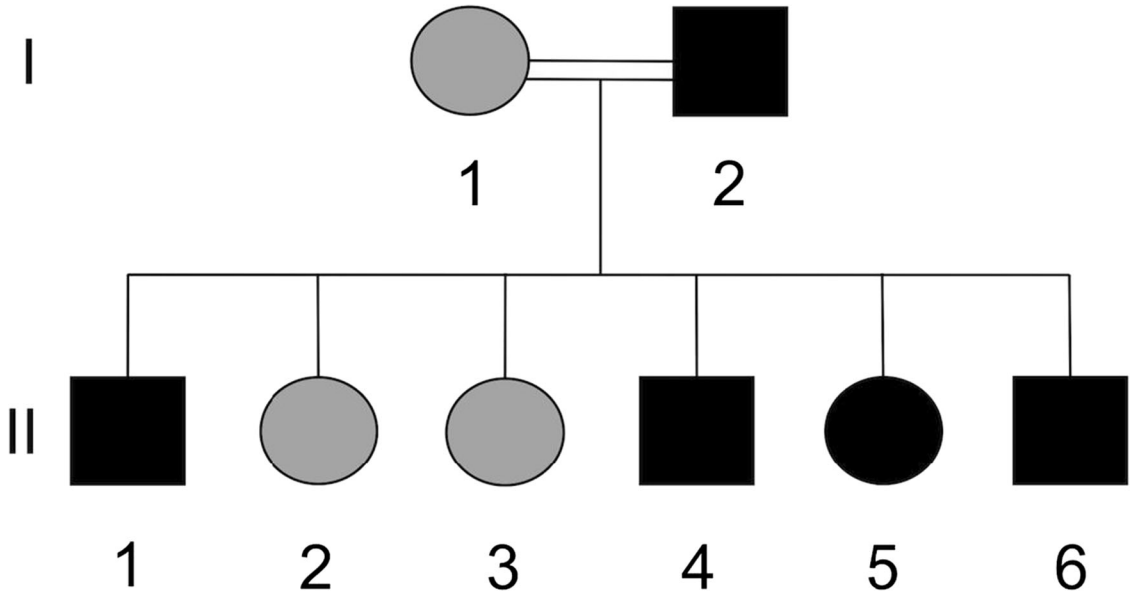

Fig. 1 Pedigree of the family with missense variants c.1604G>A (p.Gly535Asp) in *SLC24A4*. Symbols: Males are marked as squares and females as circles. Black symbols mark individuals with homozygous missense variants and severe hypomaturated amelogenesis imperfecta. Gray symbols mark individuals with heterozygous missense variants and a mild phenotype. White symbols mark clinically nonaffected individuals without a variant in SLC24A4